

Supplementary Appendix

This appendix has been provided by the authors to give readers additional information about their work.

Supplement to: Giuliano AR, Palefsky JM, Goldstone S, et al. Efficacy of quadrivalent HPV vaccine against HPV infection and disease in males. *N Engl J Med* 2011;364:401-11.

Supplemental online only materials

METHODS

Trial completion

Following demonstration of high efficacy against EGL as shown in the current analyses, and with concurrence from the Data Safety Monitoring Board (DSMB), this study was terminated to allow for unblinding and vaccination of subjects who received placebo.

Pathology panel

The independent pathology panel provided histopathologic diagnoses for endpoint identification purposes. This panel was made up of 4 experts in the field of HPV pathology. To remove medico-legal pressures, the Pathology Panel members were informed that their diagnoses would not be used for primary medical management. They were told that if their consensus diagnosis was worse than that of the Program Central Pathology Laboratory, study sites would be notified. The experts comprising the panel were Robert Kurman, M.D., Brigitte Ronnett, M.D., Mark Stoler, M.D. and Alexander Ferenczy, M.D.

Sample collection

Swabs were collected separately from the penile, scrotal, perineal/perianal and anal areas. A metal nail file was used to gently rub the penile skin. A Dacron™ swab moistened with sterile saline was used to collect cellular debris generated and then placed into a container of sample transport medium (STM)(DIGENE, Gaithersburg MD). The procedure was repeated for the scrotum and perineal/perianal regions; each swab was placed in a separate STM vial. A fourth sample from the anal canal was obtained after a sample for cytology by inserting a moistened, non lubricated Dacron™ swab a distance of 2-3 centimeters into the canal. The swab was rotated while moving it in and out several times to retrieve cells.

Milli Merck units

The milli Merck unit (mMU) is an arbitrary unit of concentration assigned for each individual HPV type. The unit is calculated based on a serially diluted reference standard for each individual HPV type in the cLIA assay. It is important to note that the mMU/ml concentration for one HPV type is not equivalent to that of another HPV type.

Table S1. Subject Accounting for the Efficacy Analysis Populations by Vaccination Group

	qHPV (N=2,032)	Placebo (N=2,033)	Total (N=4,065)	Population to Which Exclusion Category Applies	
				PPE	NRT
Number of Subjects who received at least 1 injection[†]	2,025	2,030	4,055		
Eligible for the PPE Analysis Related to:					
HPV 6/11	1,277	1,280	2,557		
HPV 16	1,327	1,308	2,635		
HPV 18	1,367	1,395	2,762		
Ineligible for the PPE Analysis Related to:					
HPV 6/11	748	750	1,498		
HPV 16	698	722	1,420		
HPV 18	658	635	1,293		
Reason for Ineligibility[‡]					
General protocol violation	83	68	151	x	
Received incorrect clinical material or dose amount	5	1	6	x	
Received non-study vaccination [§]	30	23	53	x	
Received immunosuppressives, IgG, or blood products	13	12	25	x	
With a history of immune disorder	7	9	16	x	
Diagnosed with HIV or is otherwise immunocompromised	25	15	40	x	
Received vaccine stored outside acceptable temperature range	7	9	16	x	
Subject prematurely unblinded	1	0	1	x	
Vaccination series not completed within 12 months	7	7	14	x	
Missed 2 nd and 3 rd vaccination	89	101	190	x	
Missed 3 rd vaccination	76	83	159	x	
Without Day 1 serology results within acceptable day range [¶]	5	8	13	x	x
Without Day 1 swab PCR results within acceptable day range [#]	169	161	330	x	x
Without Month 7 swab PCR results within acceptable day range [#]	244	221	465	x	
HPV 6 or 11 Positive by Serology or PCR ^{††}					
At Day 1	194	191	385	x	x
At or before Month 7	229	243	472	x	
HPV 16 Positive by Serology or PCR ^{††}					
At Day 1	120	143	263	x	x
At or before Month 7	164	192	356	x	
HPV 18 Positive by Serology or PCR ^{††}					
At Day 1	77	70	147	x	x
At or before Month 7	101	95	196	x	
[†] Subjects who did not receive at least 1 injection were excluded from all analysis populations. [‡] Subjects are counted once in each applicable exclusion category. A subject may appear in more than one category. [§] Includes live virus vaccines received within 21 days before or 14 days after study vaccination or inactivated or recombinant vaccines received within 14 days of study vaccination. Among subjects who received all 3 vaccinations. [¶] Includes subjects with a missing serum sample or missing cLIA results for ≥ 1 HPV type. [#] Includes HM subjects who are missing at least two required swab samples or PCR results for ≥ 1 HPV type and MSM subjects who are missing at least three required swab samples or PCR results for ≥ 1 HPV type. ^{††} Day 1 includes seropositivity or PCR positivity. Post-Day 1 includes PCR positivity only. Applies only to the analysis populations for the respective HPV type(s). N = Number of subjects randomized to the respective vaccination group.					

cLIA = Competitive Luminex immunoassay; HIV = Human immunodeficiency virus; HM = Heterosexual men; NRT = naïve to the relevant type; HPV = Human papillomavirus; MSM = Men having sex with men; PCR = Polymerase chain reaction; PPE = Per-Protocol efficacy; qHPV Vaccine = Quadrivalent Human Papillomavirus (Types 6, 11, 16, 18) Recombinant Vaccine.

Table S2. Baseline characteristics of subjects randomized by vaccination group

	qHPV (N = 2,032)	Placebo (N = 2,033)	Total (N = 4,065)
	n (%)	n (%)	n (%)
Sexual orientation			
Heterosexual men	1,731 (85.2)	1,732 (85.2)	3,463 (85.2)
Men having sex with men	301 (14.8)	301 (14.8)	602 (14.8)
Age (years)			
Mean (SD)	20.6 (2.0)	20.5 (2.0)	20.5 (2.0)
Median	20	20	20
Range	16 to 26	15 to 27	15 to 27
Race/Ethnicity			
Asian	201 (9.9)	205 (10.1)	406 (10.0)
Black	405 (19.9)	400 (19.7)	805 (19.8)
Hispanic American	412 (20.3)	423 (20.8)	835 (20.5)
Native American	1 (0.0)	2 (0.1)	3 (0.1)
White	719 (35.4)	712 (35.0)	1,431 (35.2)
Other	294 (14.5)	291 (14.3)	585 (14.4)
Region			
Africa	268 (13.2)	270 (13.3)	538 (13.2)
Asia-Pacific	180 (8.9)	181 (8.9)	361 (8.9)
Europe	248 (12.2)	248 (12.2)	496 (12.2)
Latin America	788 (38.8)	787 (38.7)	1,575 (38.7)
North America	548 (27.0)	547 (26.9)	1,095 (26.9)
Circumcision			
Yes	786 (38.7)	757 (37.2)	1,543 (38.0)
No	1,244 (61.2)	1,275 (62.7)	2,519 (62.0)
Lifetime number of male or female sexual partners at enrollment among non-virgins			
1	409 (20.2)	448 (22.1)	857 (21.1)
2	384 (18.9)	408 (20.1)	792 (19.5)
3	436 (21.5)	447 (22.0)	883 (21.8)
4	425 (20.9)	364 (17.9)	789 (19.4)
5	359 (17.7)	347 (17.1)	706 (17.4)
>5	2 (0.1)	6 (0.3)	8 (0.2)
Number of new male or female sexual partners in the 6 months prior to study start among non-virgins			
Unknown	2 (0.1)	0 (0.0)	2 (0.0)
0	1,149 (56.6)	1,126 (55.5)	2,275 (56.0)
1	669 (33.0)	685 (33.7)	1,354 (33.4)

2	159 (7.8)	158 (7.8)	317 (7.8)
3	30 (1.5)	38 (1.9)	68 (1.7)
4	6 (0.3)	8 (0.4)	14 (0.3)
5	2 (0.1)	3 (0.1)	5 (0.1)
>5	0 (0.0)	0 (0.0)	0 (0.0)
Percent calculated as 100*(n/N) N = number of subjects randomized; n = number of subjects with the indicated characteristic; SD = Standard deviation.			

Table S3. Analysis of efficacy in the naïve to the relevant type (NRT)* population against HPV 6/11/16/18-related (A) EGL (condyloma acuminata, PIN, and penile/perianal/perineal cancer) and (B) persistent infection and DNA detection

A. EGL

	qHPV vaccine				Placebo				Observed Efficacy (%)	95% CI
	n	Cases	Person-Years at Risk	Rate	n	Cases	Person-Years at Risk	Rate		
HPV 6/11/16/18-related EGL**	1,775	13	4,262.8	0.30	1,770	52	4,186.0	1.24	75.5	(54.3, 87.7)
By Sexual Orientation										
HM Subjects	1,501	11	3,810.8	0.29	1,497	42	3,731.8	1.13	74.4	(49.3, 88.1)
MSM Subjects	274	2	452.0	0.44	273	10	454.2	2.20	79.9	(5.7, 97.9)
HPV 6/11/16/18-related EGL by HPV type										
HPV 6-related EGL	1,603	10	3,904.8	0.26	1,607	36	3,866.3	0.93	72.5	(43.4, 87.8)
HPV 11-related EGL	1,603	1	3,919.3	0.03	1,607	16	3,879.6	0.41	93.8	(60.2, 99.9)
HPV 16-related EGL	1,674	1	4,060.0	0.02	1,649	3	3,961.7	0.08	67.5	(-305.1, 99.4)
HPV 18-related EGL	1,713	2	4,146.9	0.05	1,715	1	4,107.4	0.02	-98.1	(-11587.0, 89.7)
HPV 6/11/16/18-related EGL by EGL type										
Condyloma acuminata	1,775	10	4,268.6	0.23	1,770	48	4,187.9	1.15	79.6	(59.1, 90.8)
All PIN lesions	1,775	4	4,274.0	0.09	1,770	4	4,223.5	0.09	1.2	(-430.5, 81.6)
PIN 1	1,775	2	4,278.9	0.05	1,770	3	4,225.0	0.07	34.2	(-474.7, 94.5)
PIN 2/3	1,775	2	4,276.1	0.05	1,770	1	4,223.9	0.02	-97.6	(-11555.6, 89.7)
Penile/perianal/perineal cancer	1,775	0	4,280.9	0.00	1,770	0	4,225.4	0.00	NA	NA
<p>* This population was DNA negative to HPV 6/11/16/18/31/33/35/39/45/51/52/56/58/59 at enrollment and seronegative for HPV 6/11/16/18 at enrollment and received at least one dose of vaccine or placebo. This population approximates a population of young men prior to sexual debut. NRT case counting began after day 1.</p> <p>** Subjects are counted once in each applicable category. A subject may appear in more than one category.</p> <p>n = number of subjects who have at least one follow-up visit after month 7; Rate = rate per 100 person years at risk; CI = confidence interval; EGL = external genital lesions with a diagnosis of condyloma acuminata, PIN, or penile/perianal/perineal cancer; PIN = penile/perianal/perineal intraepithelial neoplasia.</p> <p>NRT = naïve to the relevant HPV type; this population consisted of subjects who received at least 1 dose of vaccine or placebo and were HPV-naïve (i.e., seronegative and PCR negative) to the vaccine HPV type being analyzed at day 1.</p>										

B. Persistent infection and DNA detection

	qHPV vaccine				Placebo				Observed Efficacy (%)	95% CI
	n	Cases	Person-Years at Risk	Rate	n	Cases	Person-Years at Risk	Rate		
Persistent infection*										
HPV 6/11/16/18-related persistent infection	1,669	58	3,865.2	1.50	1,664	175	3,697.2	4.73	68.3	(57.1, 76.9)
By Sexual Orientation										
HM Subjects	1,410	46	3,481.0	1.32	1,405	136	3,328.9	4.09	67.7	(54.5, 77.4)
MSM Subjects	259	12	384.2	3.12	259	39	368.3	10.59	70.5	(42.5, 85.9)
By HPV Type										
HPV 6-related persistent infection	1,513	16	3,594.0	0.45	1,510	64	3,525.1	1.82	75.5	(57.1, 86.8)
HPV 11-related persistent infection	1,513	3	3,614.9	0.08	1,510	22	3,576.4	0.62	86.5	(55.1, 97.4)
HPV 16-related persistent infection	1,578	29	3,703.2	0.78	1,545	76	3,567.4	2.13	63.2	(42.9, 76.9)
HPV 18-related persistent infection	1,610	13	3,799.6	0.34	1,614	42	3,744.0	1.12	69.5	(42.1, 85.0)
DNA detection										
HPV 6/11/16/18-related DNA detection	1,669	253	3,669.7	6.89	1,664	372	3,547.9	10.49	34.2	(22.7, 44.2)
By Sexual Orientation										
HM Subjects	1,410	193	3,313.1	5.83	1,405	299	3,192.2	9.37	37.8	(25.2, 48.4)
MSM Subjects	259	60	356.7	16.82	259	73	355.7	20.52	18.0	(-16.9, 42.7)
By HPV Type										
HPV 6-related DNA detection	1,513	88	3,533.2	2.49	1,510	161	3,457.8	4.66	46.5	(30.2, 59.2)
HPV 11-related DNA detection	1,513	27	3,593.6	0.75	1,510	54	3,554.3	1.52	50.5	(20.1, 70.0)
HPV 16-related DNA detection	1,578	119	3,606.9	3.30	1,545	164	3,508.5	4.67	29.4	(10.1, 44.7)
HPV 18-related DNA detection	1,610	59	3,752.5	1.57	1,614	106	3,704.8	2.86	45.0	(23.7, 60.7)
<p>*Persistent infection was defined as detection of the same HPV type (6, 11, 16 or 18) in an anogenital swab or biopsy specimen collected on ≥ 2 consecutive visits ≥ 6 (± 1) months apart. Individuals with detection of HPV 6, 11, 16, or 18 DNA at ≥ 1 visit contributed to the DNA detection endpoint.</p> <p>n = number of subjects who have at least one follow-up visit after day 1; Rate = rate per 100 person years at risk; CI = confidence interval.</p>										

Table S4. Analysis of efficacy against HPV 6/11/16/18-related persistent infection and DNA detection in the per-protocol efficacy (PPE) population

	qHPV vaccine				Placebo				Observed Efficacy (%)	CI [†]	P-value [‡]
	n	Cases	Person-Years at Risk	Rate	n	Cases	Person-Years at Risk	Rate			
Persistent infection*											
HPV 6/11/16/18-related persistent infection	1,390	15	2,549.4	0.59	1,400	101	2,469.3	4.09	85.6	(73.4, 92.9)	< 0.001
By Sexual Orientation											
HM Subjects	1,196	14	2,356.5	0.59	1,195	83	2,274.0	3.65	83.7	(71.1, 91.5)	
MSM Subjects	194	1	192.9	0.52	205	18	195.3	9.22	94.4	(64.4, 99.9)	
By HPV Type											
HPV 6-related persistent infection	1,239	4	2,320.2	0.17	1,238	33	2,296.6	1.44	88.0	(66.3, 96.9)	
HPV 11-related persistent infection	1,239	1	2,322.6	0.04	1,238	15	2,315.1	0.65	93.4	(56.8, 99.8)	
HPV 16-related persistent infection	1,290	9	2,382.4	0.38	1,264	41	2,312.9	1.77	78.7	(55.5, 90.9)	
HPV 18-related persistent infection	1,327	1	2,461.9	0.04	1,347	25	2,453.5	1.02	96.0	(75.6, 99.9)	
DNA detection											
HPV 6/11/16/18-related DNA detection	1,390	136	2,455.3	5.54	1,400	241	2,404.1	10.02	44.7	(31.5, 55.6)	< 0.001
By Sexual Orientation											
HM Subjects	1,196	115	2,270.7	5.06	1,195	199	2,215.1	8.98	43.6	(28.7, 55.6)	
MSM Subjects	194	21	184.6	11.38	205	42	189.0	22.22	48.8	(11.6, 71.2)	
By HPV Type											
HPV 6-related DNA detection	1,239	51	2,292.4	2.22	1,238	99	2,267.7	4.37	49.0	(27.9, 64.4)	
HPV 11-related DNA detection	1,239	16	2,311.7	0.69	1,238	37	2,300.5	1.61	57.0	(20.7, 77.6)	
HPV 16-related DNA detection	1,290	62	2,337.7	2.65	1,264	103	2,287.8	4.50	41.1	(18.5, 57.7)	
HPV 18-related DNA detection	1,327	25	2,441.3	1.02	1,347	66	2,440.6	2.70	62.1	(39.2, 77.1)	
<p>*Persistent infection was defined as detection of the same HPV type (6, 11, 16 or 18) in an anogenital swab or biopsy specimen collected on ≥ 2 consecutive visits ≥ 6 (± 1) months apart. Individuals with detection of HPV 6, 11, 16, or 18 DNA at ≥ 1 visit contributed to the DNA detection endpoint.</p> <p>[†] A 97.5% CI is reported for the HPV 6/11/16/18-related persistent infection endpoint. The CI reported for the HPV 6/11/16/18-related persistent infection endpoint differs from the other analyses due to the Hochberg multiplicity adjustment applied.</p> <p>[‡] A p-value < 0.025 (one-sided) corresponds to a lower bound of the confidence interval for vaccine efficacy greater than 20% and supports the conclusion that the vaccine is efficacious against the given endpoint. The Hochberg multiplicity adjustment has been applied to the p-value reported.</p> <p>n = number of subjects who have at least one follow-up visit after month 7; Rate = rate per 100 person years at risk; CI = confidence interval.</p>											

Table S5. Breakdown of injection-site and systemic adverse experiences reported following any vaccination (incidence $\geq 1\%$ in one or more vaccination group).

	qHPV vaccine				Placebo			
	All AE		Vaccine related AE [‡]		All AE		Vaccine related AE [‡]	
	n	(%)	n	(%)	n	(%)	n	(%)
Subjects in analysis population	2,020				2,029			
Subjects with follow up	1,945				1,950			
Injection-site related*	1166	(59.9)			1046	(53.6)		
Erythema	304	(15.6)			275	(14.1)		
Pain**	1113	(57.2)			991	(50.8)		
Pruritus	22	(1.1)			24	(1.2)		
Swelling	219	(11.3)			187	(9.6)		
Systemic (by system organ class)[†]	615	(31.6)	274	(14.1)	613	(31.4)	284	(14.6)
Gastrointestinal disorders	125	(6.4)	35	(1.8)	120	(6.2)	33	(1.7)
Abdominal pain - upper	19	(1.0)	5	(0.3)	23	(1.2)	7	(0.4)
Diarrhea	40	(2.1)	10	(0.5)	36	(1.8)	13	(0.7)
Nausea	27	(1.4)	16	(0.8)	16	(0.8)	7	(0.4)
General disorders	160	(8.2)	109	(5.6)	169	(8.7)	122	(6.3)
Fatigue	13	(0.7)	6	(0.3)	19	(1.0)	15	(0.8)
Pyrexia	118	(6.1)	91	(4.7)	125	(6.4)	98	(5.0)
Infections and infestations	182	(9.4)	18	(0.9)	187	(9.6)	20	(1.0)
Influenza	42	(2.2)	9	(0.5)	44	(2.3)	7	(0.4)
Nasopharyngitis	44	(2.3)	3	(0.2)	50	(2.6)	5	(0.3)
Pharyngitis	22	(1.1)	1	(0.1)	20	(1.0)	0	(0.0)
Upper respiratory tract infection	27	(1.4)	3	(0.2)	20	(1.0)	4	(0.2)
Injury, poisoning and procedural complications	30	(1.5)	0	(0.0)	24	(1.2)	0	(0.0)
Musculoskeletal disorders	61	(3.1)	21	(1.1)	50	(2.6)	15	(0.8)
Nervous system disorders	207	(10.6)	121	(6.2)	231	(11.8)	138	(7.1)
Dizziness	19	(1.0)	12	(0.6)	18	(0.9)	14	(0.7)
Headache	179	(9.2)	107	(5.5)	207	(10.6)	119	(6.1)
Respiratory disorders	70	(3.6)	25	(1.3)	68	(3.5)	8	(0.4)
Pharyngolaryngeal pain	38	(2.0)	14	(0.7)	37	(1.9)	2	(0.1)
Skin and subcutaneous disorders	26	(1.3)	10	(0.5)	31	(1.6)	14	(0.7)
<p>*Days 1-5 following vaccination; all injection site adverse experiences are considered vaccine-related.</p> <p>**P value for risk difference between vaccine and placebo groups = 0.001</p> <p>[†]Days 1-15 following vaccination.</p> <p>[‡] Determined by the investigator to be possibly, probably, or definitely related to the vaccine.</p>								

Although a subject may have had two or more adverse experiences, the subject is counted only once within a category.
Percentages are calculated based on the number of subjects with follow-up.

Figure S1. Subject accounting in Protocol 020

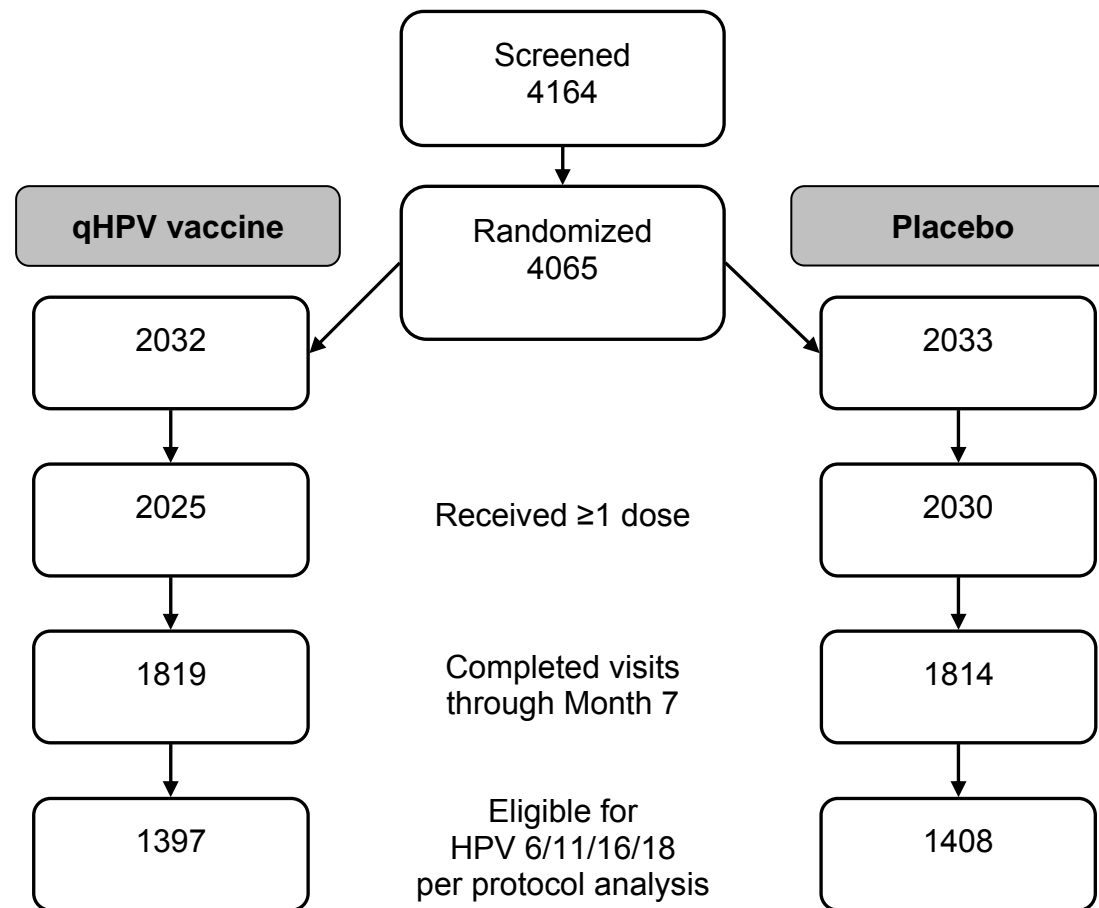
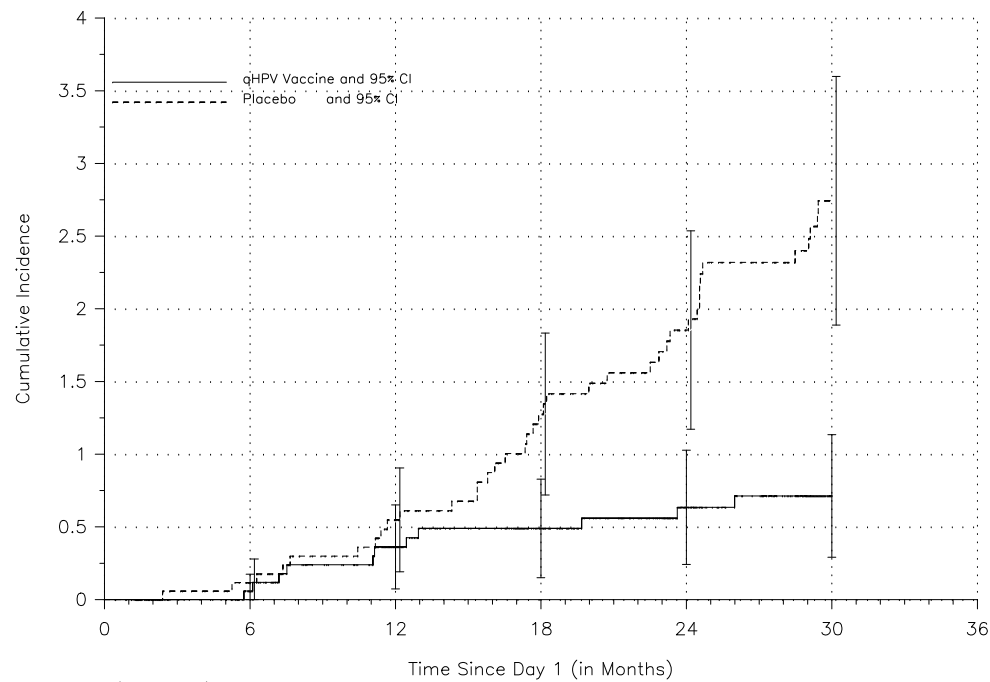


Figure S2. Analysis of time to 6/11/16/18-related EGL in the NRT population



Number of Subjects at Risk						
qHPV Vaccine						
Placebo	1,775	1,686	1,582	1,455	1,311	1,090
	1,770	1,682	1,569	1,422	1,275	1,041

EGL = external genital lesions with a diagnosis of condyloma acuminata, penile/perianal/perineal intraepithelial neoplasia, or penile/perianal/perineal cancer; Cumulative incidence = Cumulative incidence per 100 person years; qHPV vaccine = quadrivalent human papillomavirus (types 6, 11, 16, 18) recombinant vaccine.